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Figures

Fig 1.

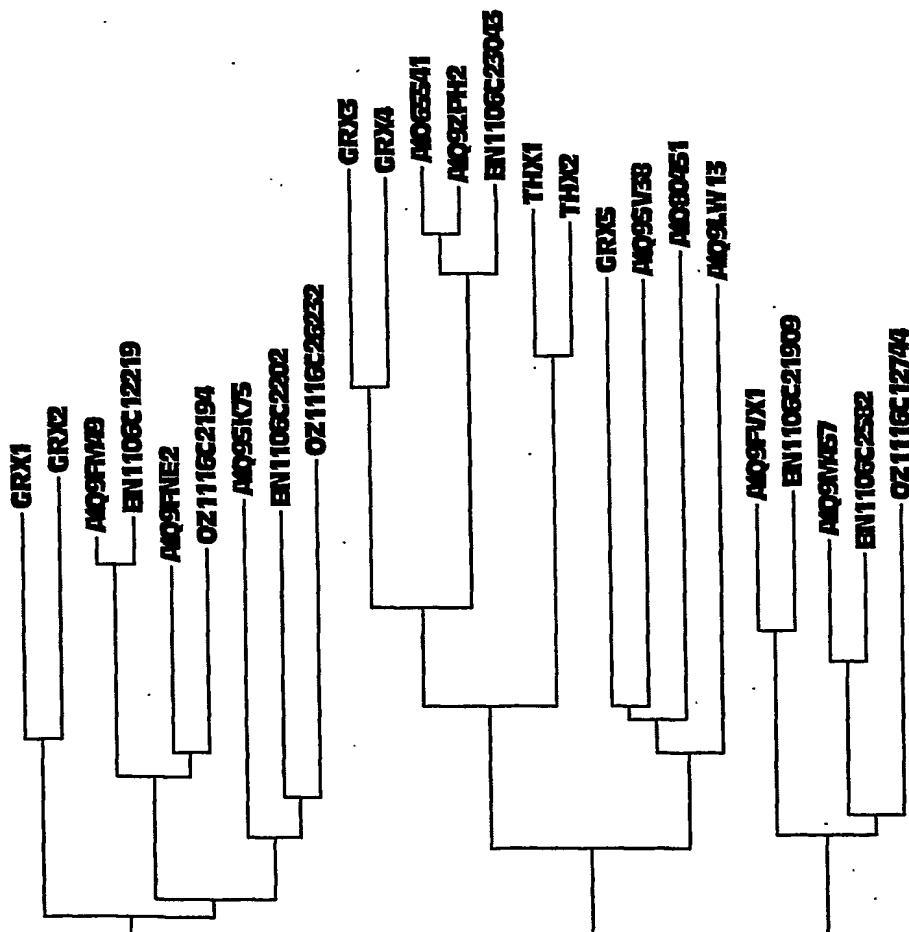


Fig. 2

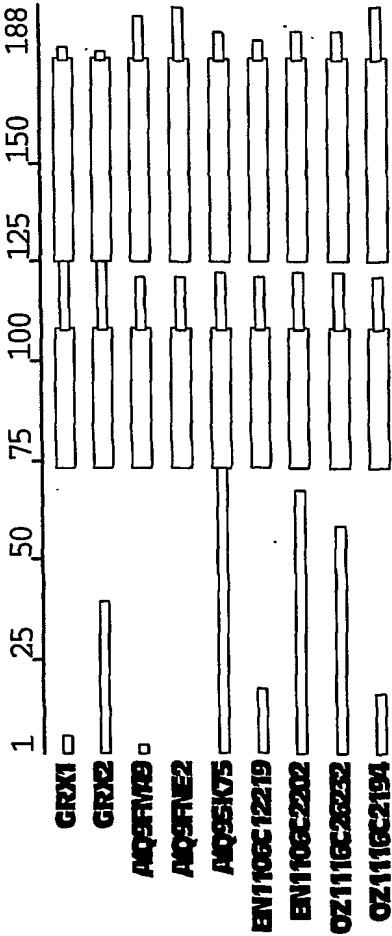
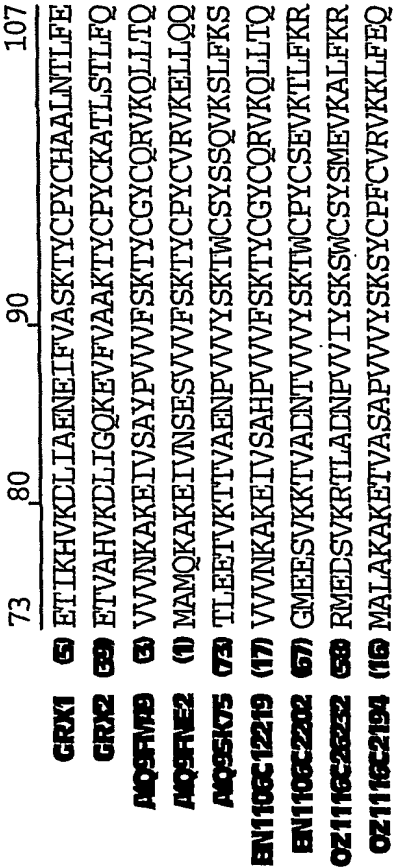


Fig. 3



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Fig. 4

	125	130	140	150	160	175
GRX1 (57)	KEGADIQAALYEINGQRTVPNIYINGKHHIGGNDLQELRETGELEELLEPI					
GRX2 (91)	SNGSEIQDALEEISGQKTVPNVYINGKHHIGGNSDLETLLKKNCKLAELKPV					
ATQ9FM49 (51)	SDGGEIQSALSSEWTGQTTVPNVFIKGNHHIGGCDRVMETNKQKLVPLLTEA					
ATQ9FNE2 (49)	SDGSQIQSGLAEWTGQRTVPNVFIKGNHHIGGCDATSNLHKDGLVPLLTEA					
ATQ95K75 (122)	SEGSQIQNVLEKITGQYTVPNVFIKGNHHIGGCDTLQHLNKGLEAIIAAEA					
BN1106C12219 (65)	SDGGEIQSALSSEWTGQSTVPNVFIKGNHHIGGCDRVMESNKQKLVPLLTEA					
BN1106C2202 (116)	PQGTQLQKVLETLTGQRTVPNVFVGGKHHIGGCTDTVNLNRKGLELMLAAEA					
OZ1116C26232 (107)	AQGPQLQKVLERLTGQSTVPNVFIKGNHHIGGCTDTVKLHRKGELATMLSEL					
OZ1116C2194 (64)	SDGSELOSALEWTGQRTVPNVFINGKHHIGGCDDTLALNNEGKLVPLLTEA					

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Fig. 5

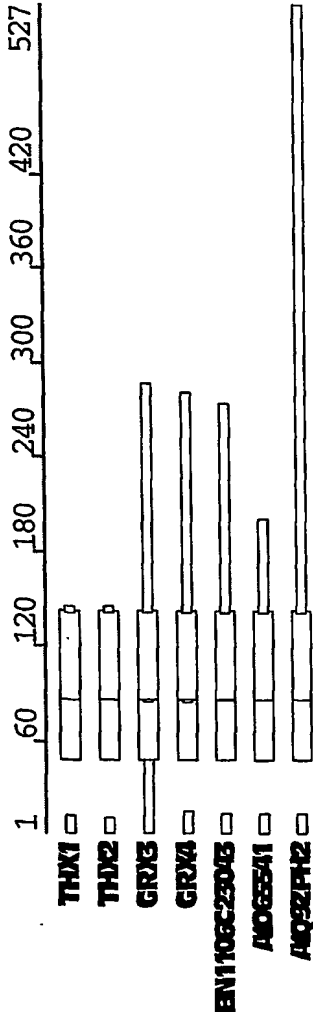
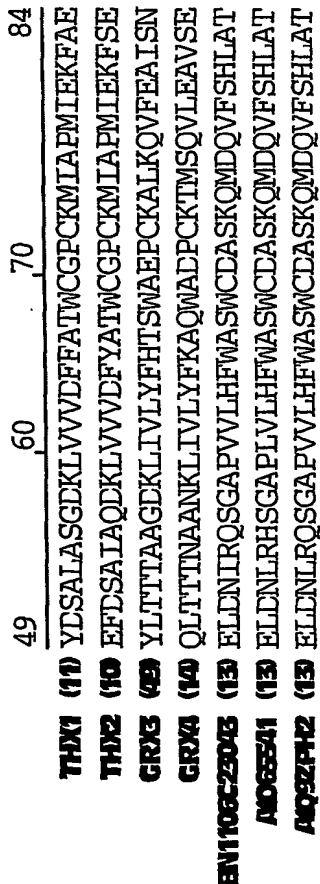


Fig. 6



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Fig. 7

	87	100	110	120	139																																																
THX1 (47)	Q	Y	S	D	A	A	F	Y	K	L	D	V	D	E	V	S	D	V	A	Q	K	A	E	V	S	S	M	P	T	L	I	F	Y	K	G	G	K	E	V	T	R	V	V	G	A	N	P	A	A	I	K	Q	A
THX2 (46)	Q	Y	P	Q	A	D	F	Y	K	L	D	V	D	E	L	G	D	V	A	Q	K	N	E	V	S	A	M	P	T	L	L	L	F	K	N	G	K	E	V	A	K	V	V	G	A	N	P	A	A	I	K	Q	A
GRX3 (87)	S	N	S	N	V	S	F	L	S	I	D	A	D	E	N	S	E	I	S	E	L	F	E	I	S	A	V	P	F	I	I	H	K	G	T	I	L	K	E	L	S	G	A	D	P	K	E	Y	V	S	L		
GRX4 (52)	R	Q	E	D	V	R	F	L	S	I	D	A	D	E	H	P	E	I	S	D	L	F	E	I	A	A	V	P	F	V	F	I	Q	N	G	T	I	V	K	E	I	S	A	A	D	P	K	E	F	V	K	S	
BN1106C23043 (49)	D	F	P	R	A	H	F	F	R	V	E	A	E	E	H	P	E	I	S	E	A	Y	S	V	S	A	V	P	F	V	F	F	K	D	G	K	A	V	D	T	L	E	G	A	D	P	S	S	L	A	N	K	
A1065541 (49)	D	F	P	R	A	H	F	F	R	V	E	A	E	E	H	P	E	I	S	E	A	Y	S	V	A	L	V	P	F	V	F	F	K	D	G	K	T	V	D	T	L	E	G	A	D	P	S	S	L	A	N	K	
A1092PH2 (49)	D	F	P	R	A	H	F	F	R	V	E	A	E	E	H	P	E	I	S	E	A	Y	S	V	A	A	V	P	F	V	F	F	K	D	G	K	T	V	D	T	L	E	G	A	D	P	S	S	L	A	N	K	

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Fig. 8

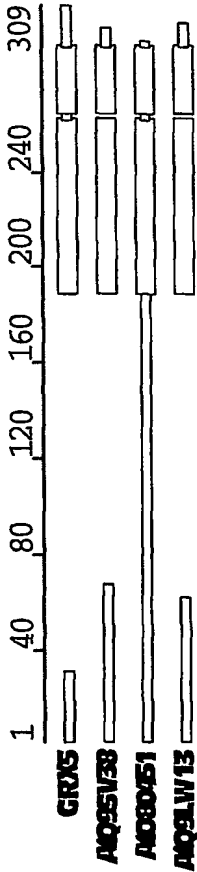


Fig. 9

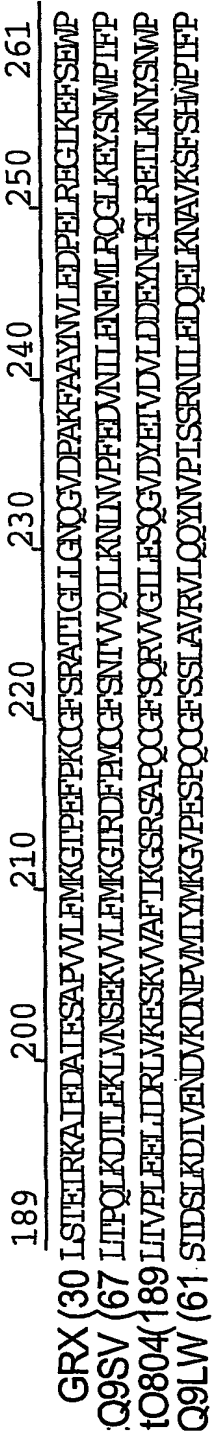
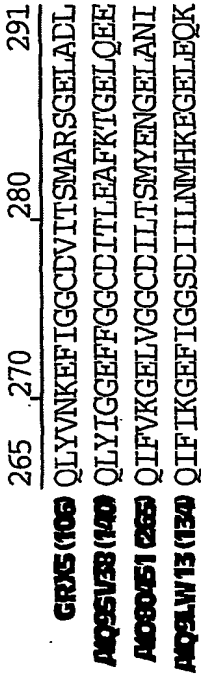


Fig. 10



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Fig. 11

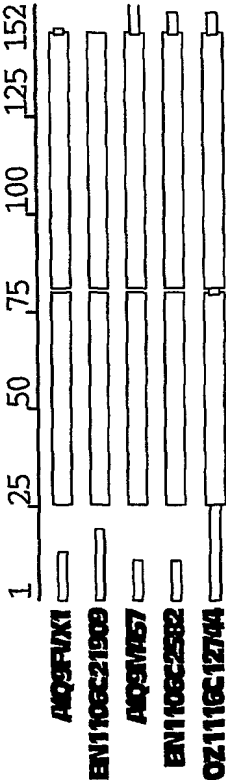


Fig. 12

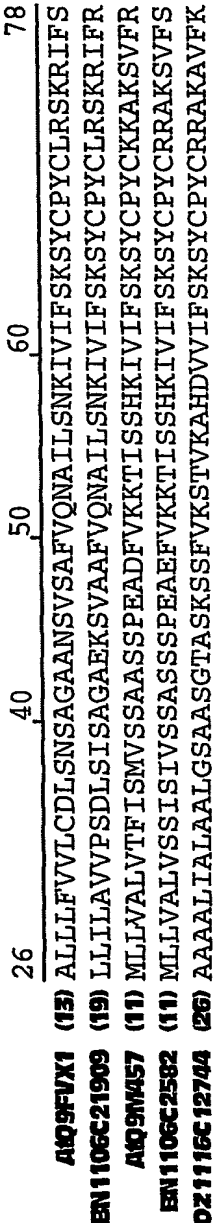


Fig. 13

